Figure 1:

	SCS0007	MWAGLLLRAACVALLLPGAPARGYTGRKPPGHFAAERRRLGPHV
	FLJ32009	MWAGLLLRAACVALLLPGAPARGYTGRKPPGHFAAERRRLGPHV
_	HCP50656.1	RGAGAEPSPGDPPPPVMWAGLLLRAACVALLLPGAPARGYTGRKPPGHFAAE RRRLGPHV
5		
	scs0007	CLSGFGSGCCPGWAPSMGGGHCTLPLCSFGCGSGICIAPNVCSCQDGEQGATCPETHGPC
	FLJ32009	CLSGFGSGCCPGWAPSMGGGHCTLPLCSFGCGSGICIAPNVCSCQDGEQGAT CPETHGPC
	HCP50656.1	CLSGFGSGCCPGWAPSMGGGHCTLPLCSFGCGSGICIAPNVCSCQDGEQGATCPETHGPC
10		***************************************
	scs0007	GEYGCDLTCNHGGCQEVARVCPVGFSMTETAVGIRCTDIDECVTSSCEGHCV NTEGGFVC
	FLJ32009	GEVCCDLTCNHGGCOEVARVCPVGFSMTEAAVGIRCTDIDECVTSSCEGHCVNTEGGEVC .
	HCP50656.1	GEYGCDLTCNHGGCQEVARVCPVGFSMTETAVGIRCTDIDECVTSSCEGHCVNTEGGFVC
15		******
	scs0007	ECGPGMQLSADRHSCQDTDECLGTPCQQRCKNSIGSYKCSCRTGFHLHGNRHSCVDVNEC
	FLJ32009	ECGPGMOLSADRHSCODTDECLGTPCOORCKNSIGSYKCSCRTGFHLHGNRHSCVDVNEC
	HCP50656.1	ECGPGMQLSADRHSCQDTDECLGTPCQQRCKNSIGSYKCSCRTGFHLHGNRHS CVDVNEC
20		****
	scs0007	RRPLERRVCHHSCHNTVGSFLCTCRPGFRLRADRVSCEG
	FLJ32009	RRPLERRVCHHSCHNTVGSFLCTCRPGFRLRADRVSCEAFPKAVLAPSAILQP RQHPSKM
	HCP50656.1	RRPLERRVCHHSCHNTVGSFLCTCRPGFRLRADRVSCEG
25		***********
	scs0007	
	FLJ32009	LLLLPEAGRPALSPGHSPPSGAPGPPAGVRTTRLPSPTPRLPTSSPSAPVWLLSTLLATP
	HCP50656.1	
30		****
	scs0007	WEPCMNQGVAGQSLG
	FLJ32009	VPTASLLGNLRPPSLLQGEVMGTPSSPRGPESPRLAAGPSPCWHLGAMHESRSRWTEPGC
	HCP50656.1	** * *
35		•
	SCS0007	
	FLJ32009	SQCWCEDGKVTCEKVRCEAACSHPIPSRDGGCCPSCTGCFHSGVVRAEGDVFSP PNENCT
	HCP50656.1	** * * * *
40		
	scs0007	VLLARVRPPIKTDCCTCVPVRCYFHGRWYADGAVFSGGG DECTTC
	FLJ32009	VCVCLAGNVSCISPECPSGPCOTPPOTDCCTCVPVRCYFHGRWYADGAVFSGGGDECTTC
	HCP50656.1	VLLARVRPPIKTDCCTCVPVRCYFHGRWYADGAVFSGGDECTTC
45		
	scs0007	VCQNGEVECSFMPCPELACPREEWRLGPGQCCFTCQEPTPSTGCSLDDNGVEFPIGQIWS
	FLJ32009	VCONCEVECSEMPCPELACPREEWRLGPGOCCFTCQEPTPSTGCSLDDNGVEFP1GQ1WS
	HCP50656.1	VCQNGEVECSFMPCPELACPREEWRLGPGQCCFTCQEPTPSTGCSLDDNGVEFPI GQIWS
50		
	scs0007	PGDPCELCICQADGSVSCKRTDCVDSCPHPIRIPGQCCPDCSAAGAQRMLSLAGCTYTGR
	FLJ32009	PGDPCELCICOADGSVSCKRTDCVDSCPHPIRIPGQCCPDCSAGCTYTGR
	HCP50656.1	PGDPCELCICQADGSVSCKRTDCVDSCPHPIRIPGQCCPDCSAAGAQRMLSLAGCTYTGR
55		有有有有有有有有有有有有有有有有有有有有有有有有有有有有有有有有有有有有
	scs0007	IFYNNETFPSVLDPCLSCICLLGSVACSPVDCPITCTYPFHPDGECCPVCRDCNY EGRKV
	FLJ32009	TEVNNETEPSVLDPCLSCICLLGSVACSPVDCPITCTYPFHPDGECCPVCRDCNYEGRKV
	HCP50656.1	IFYNNETFPSVLDPCLSCICLLGSVACSPVDCPITCTYPFHPDGECCPVCRDCNYEGRKV
60		**********

 ${\tt ANGQVFTLDDEPCTRCTCQLGEVSCEKVPCQRACADPALLPGDCCSSCPDSLSPLEEKQG}$ SCS0007 ANGQVFTLDDEPCTRCTCQLGEVSCEKVPCQRACADPALLPGDCCSSCPDSLSPLEEKQG FLJ32009 ANGQVFTLDDEPCTRCTCQLGEVSCEKVPCQRACADPAL LPGDCCSSCPDSPQPLQ----5 HCP50656.1 ********** ${\tt LSPHGNVAFSKAGRSLHGDTEAPVNCSSCPGPPTASPSRPVLHLLQLLLRTNLMKTQTLP}$ SCS0007 LSPHGNVAFSKAGRSLHGDTEAPVNCSSCPGPPTASPSR PVLHLLQLLLRTNLMKTQTLP FLJ32009 -- PPPTLAPSSPQWGLLGGKSPP ---- CCLG------FPMLHCSFTAYLWEGGKRDPKP 10 HCP50656.1 TSPAGAHGPHSLALGLTATFPGEPGASPRLSPGPSTPPG APTLPLASPGAPQPPPVTPER scs0007 TSPAGAHGPHSLALGLTATFPGEPGASPRLSPGPSTPPGAPTLPLASPGAPQPPPVTPER FLJ32009 TGGKCYFHGRWYTDGAVFSGGGDDCTTCVCQNGEVECSFTPCPELECPRE -----EW 15 HCP50656.1 . . * . *. .. SFSASGAQIVSRWPPLPGTLLTEASALSMMDPSPSKTPITLLGPRVLSP -TTSRLSTALA SFSASGAQIVSRWPPLPGTLLTEASALSMMDPSPSKTPITLLGPRVLSP -TTSRLSTALA scs0007 FLJ32009 LLGPGQCCFTCREP -----TPTTGCSLDD-----NGVEFPIGQIWSPGDPCRRLCELQ HCP50656.1 20 ${\tt ATTHPGPQQPPVGASRGEESTMLSRFPHAALLIHRLPVGRWET}$ SCS0007 ATTHPGPQQPPVGASRGEESTM -----FLJ32009 EDRLRGLLPAPHPDSWTVLS -----25 HCP50 656.1

* *

Figure 2:

	Sequence	Annotation
	AK056571	weakly similar to FIBULIN-1, ISOFORM A Precursor
	XP_166226.	similar to cysteine-rich motor neuron 1; cysteine-rich repeat-containing protein
5		S52 precursor; von Willebrand factor type C domain.
10	SCS0007 AK056571 XP_166226.	MWAGLLLRAA CVALLLPGAP ARGYTGRKPP GHFAAERRRL GPHVCLSGFG MWAGLLLRAA CVALLLPGAP ARGYTGRKPP GHFAAERRRL GPHVCLSGFG
15	SCS0007 AK056571 XP_166226.	SGCCPGWAPS MGGGHCTLPL CSFGCGSGIC IAPNVCSCQD GEQGATCPET SGCCPGWAPS MGGGHCTLPL CSFGCGSGIC IAPNVCSCQD GEQGATCPETPA LSPGHS
20	SCS0007 AK056571 XP_166226.	HGPCGEYGCD LTCNHGGCQE VARVCPVGFS MTETAVGIRC TDIDECVTSS HGPCGEYGCD LTCNHGGCQE VARVCPVGFS MTEAAVGIRC TDIDECVTSS GAP
25	SCS0007 AK056571 XP_166226.	CEGHCVNTEG GFVCECGPGM QLSADRHSCQ DTDECLGTPC QQRCKNSIGS CEGHCVNTEG GFVCECGPGM QLSADRHSCQ DTDECLGTPC QQRCKNSIGSRLPSPT PRLPTSSP
30	SCS0007 AK056571 XP_166226.	YKCSCRTGFH LHGNRHSCVD VNECR RPLER RVCHHSCHNT VGSFLCTCRP YKCSCRTGFH LHGNRHSCVD VNECRRPLER RVCHHSCHNT VGSFLCTCRPSAPVWL LSTLLATFVP TASLLGNLRP
35	SCS0007 AK056571 XP_166226.	GFRLRADRVS CEG
40	SCS0007 AK056571 XP_166226.	PDWQQGP
45	SCS0007 AK056571 XP_166226.	LGNLRPPSLL QGEVMGTPSS PRGPESPRLA AGPSPCWHLG AMHESRSRWTSPCWHLG AMHESRSRWT
50	SCS0007 AK056571 XP_166226.	EPGCSQCWCE DGKVTCEKVR CEAACSHPIP SRDGGCCPSC TGCFHSGVVR EPGCSQCWCE DGKVTCEKVR CEAACSHPIP SR DGGCCPSC TGCFHSGVVR
55	SCS0007 AK056571 XP_166226.	AEGDVFSPPN ENCTVCVCLA GNVSCISPEC PSGPCQTPPQ TDCCTCVPVR AEGDVFSPPN ENCTVCVCLA GNVSCISPEC PSGPCQTPPQ TDCCTCVPVR

5	SCS0007 AK056571 XP_166226.	CYFHGRWYAD GAVFSGGGDE CTTCVCQNGE VECSFMPCPE LACPREEWRL CYFHGRWYAD GAVFSGGGDE CTTCVCQNGE VECSFMPCPE LACPREEWRL CYFHGRWYAD GAVFSGGGDE CTTCVCQNGE VECSFMPCPE LACPREEWRL
3		
	SCS0007 AK056571	GPGQCCFTCQ EPTPSTGCSL DDNGVEFPIG QIWS PGDPCE LCICQADGSV GPGQCCFTCQ EPTPSTGCSL DDNGVEFPIG QIWSPGDPCE LCICQADGSV GPGQCCFTCQ EPTPSTGCSL DDNGVEFPIG QIWSPGDPCE LCICQADGSV
10	XP_166226.	Gragectica Principal Principal
10		
	SCS0007 AK056571 XP_166226.	SCKRTDCVDS CPHPIRIPGQ CCPDCSAAGA QRMLSLAGCT YTGRIFYNNE
	AK056571	SCK RTDCVDS CPHPIRIPGQ CCPDCSAG CT YTGRIFYNNE
	XP_166226.	SCKRTDCVDS CPHPIRIPGQ CCPDCSAGCT YTGRIFYNNE
15		
	ecennn7	TFPSVLDPCL SCICLLGSVA CSPVDCPITC TYPFHPDGEC CPVCRDCNYE TFPSVLDPCL SCICLLGSVA CSPVDCPITC TYPFHPDGEC CPVCRDCNYE TFPSVLDPCL SCICLLGSVA CSPVDCPITC TYPFHPDGEC CPVCRDCNYE
	AK056571	TFPSVLDPCL SCICLLGSVA CSPVDCPITC TYPFHPDG EC CPVCRDCNYE
	XP 166226.	TFPSVLDPCL SCICLLGSVA CSPVDCPITC TYPFHPDGEC CPVCRDCNYE
20		
		GRKVANGQVF TLDDEPCTRC TCQLGEVSCE KVPCQRACAD PALLPGDCCS
	scs0007	
	AK056571	GRKVANGQVF TEDDEPCTRC TCQLGEVSCE KVPCQRACAD PALLPGDCCS
25	XP_166226.	GRAVANG OVE THE BELLEVIE TO SEE THE SECOND S
23		
	SCS0007	SCPDSLSPLE EKQGLSPHGN VAFSKAGRSL HGDTEAPVNC SSCPGPPTAS
	SCS0007 AK056571	SCPDSLSPLE EKQGLSPHGN VAFSKAGRSL HGDTEAPVNC SSCPGPPTAS
	XP_166226.	SCPDSLSPLE EKQGLSPHGN VAFSKAGRSL HGDTEAPVNC S SCPGPPTAS
30		
	0000007	PSRPVLHLLQ LLLRTNLMKT QTLPTSPAGA HGPHSLALGL TATFPGEPGA PSRPVLHLLQ LLLRTNLMKT QTLPTSPAGA HGPHSLALGL TATFPGEPGA PSRPVLHLLQ LLLRTNLMKT QTLPTSPAGA HGPHSLALGL TATFPGEPGA
	SCS0007 NK056571	PSRPVLHILO LLLRTNLMKT OTLPTSPAGA HGPHSLALGL TATFPGEPGA
	XP 166226.	PSRPVLHLLQ LLLRTNLMKT QTLPTSPAGA HGPHSLALGL TATFPGEPGA
35		
		TOTAL DESIGNATION OF THE PROPERTY OF THE PROPE
	SCS0007	SPRLSPGPS T PPGAPTLPLA SPGAPQPPPV TPERSFSASG AQIVSRWPPL SPRLSPGPST PPGAPTLPLA SPGAPQPPPV TPERSFSASG AQIVSRWPPL
	AK056571 XP_166226.	SPRISPGPST PPGAPTIPLA SPGAPQPPPV TPERSFSASG AQIVSRWPPL
40	XP_166226.	SPRESFORM FEGRETHER STORINGSTEEL BETTER STORINGS
40		
	scs0007	PGTLLTEASA LSMMDPSPSK TPITLLGPRV LSPTTSRLST ALA ATTHPGP PGTLLTEASA LSMMDPSPSK TPITLLGPRV LSPTTSRLST ALAATTHPGP
	AK056571	PGTLLTEASA LSMMDPSPSK TPITLLGPRV LSPTTSRLST ALAATTHPGP
	XP_166226.	PGTLLTEASA LSMMDPSPSK TPITLLGPRV LSPTTSRLST ALAATTHPGP
45		
	200007	CORRUCASES FESTMISRED HAALLIHRLP VGRWET
	SCSUUU/ 3K056571	OOPPVGASRG EESTM
	XP 166226.	QQPPVGASRG EESTMLSRFP HAALLIHRLP VGRWET QQPPVGASRG EESTM
50	W-700550.	
50		